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(75) Inventors/Applicants (for US only): COLLEN, Désiré, José [BE/GB]; 28 Collingham Gardens, London SW5 0HN (GB). NAGAI, Nubuo [JP/BE]; Benedenstraat 60, B-3000 Leuven (BE). LAROCHE, Yves [BE/BE]; Bémelstraat 115, B-1150 Brussel (BE).

(21) International Application Number: PCT/BE01/00217 (74) Agents: BIRD, William, E. et al.; Bird Goën & Co, Vilvoordsebaan 92, B-3020 Winksele (BE).

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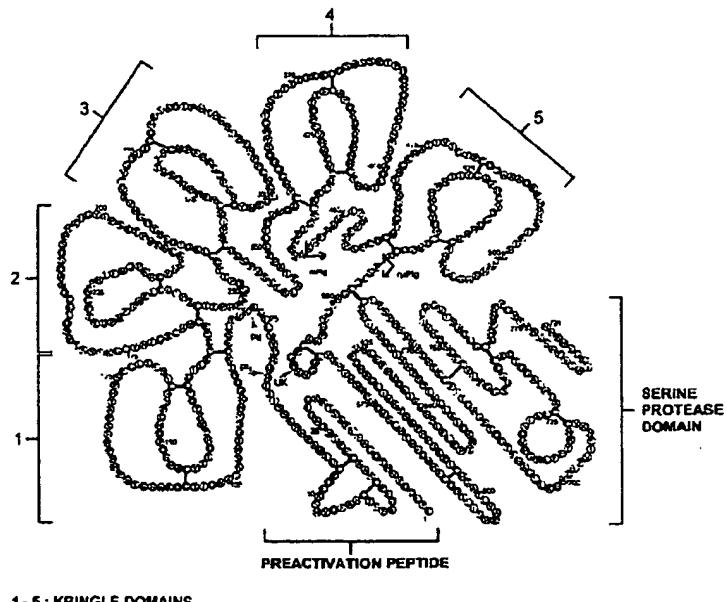
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(71) Applicant (for all designated States except US): THROMB-X NV [BE/BE]; Leopold I straat 1bus 21, B-3000 Leuven (BE). (84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW),

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(54) Title: A YEAST EXPRESSION VECTOR AND A METHOD OF MAKING A RECOMBINANT PROTEIN BY EXPRESSION IN A YEAST CELL



1-5: KRINGLE DOMAINS

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(57) Abstract: Vectors for the expression in yeast of mammalian plasminogen derivatives such as microplasminogen and miniplasminogen are presented. Methods for expression of these proteins in a methylotrophic yeast expression system are disclosed as well as the activation and stabilisation of the recombinant proteins. The proteins of this invention are used in the treatment of focal cerebral ischemic infarction and other thrombotic diseases.



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A YEAST EXPRESSION VECTOR AND A METHOD OF MAKING A RECOMBINANT PROTEIN BY EXPRESSION IN A YEAST CELL.

This invention relates to the treatment and prevention of thrombotic disorders. More specifically, the present invention relates to the high yield 5 production, via recombinant DNA technology, of derivatives of mammalian plasminogen, their purification and stabilization, and to the use of the corresponding activated and stabilized plasmin derivatives for the treatment of focal cerebral ischemic infarction (ischemic stroke) or arterial thrombotic diseases such as peripheral arterial occlusive disease or acute myocardial infarction.

10 **BACKGROUND OF THE INVENTION**

Mammalian blood contains an enzymatic system, called the fibrinolytic or plasminogen system, which plays a role in various biological phenomena such as reproduction, embryogenesis, cell invasion, angiogenesis and brain function. In addition, this system participates in thrombosis, atherosclerosis, neoplasia, 15 metastasis and chronic inflammatory disorders. The fibrinolytic system contains plasminogen, which by the action of plasminogen activators is converted to the active enzyme plasmin, which in turn digests fibrin to soluble degradation products. Two physiological plasminogen activators, respectively called tissue-type (t-PA) and urokinase-type (u-PA), have been identified. Inhibition of the 20 fibrinolytic system may occur either at the level of plasminogen activators, by means of specific plasminogen activator inhibitors (PAI), or at the level of plasmin, mainly by means of α_2 -antiplasmin.

A number of substances are involved in clot formation and lysis. Plasminogen and plasmin are two of the primary substances involved in lysis. 25 Plasminogen, a protein composed of 791 amino-acids that circulates in plasma at a concentration of about 200 μ g/ml, is the zymogen form of a fibrinolytic enzyme, plasmin, which has broad substrate specificity and is ultimately responsible for degrading blood clots. For the most part, fibrin proteolysis is mediated by the generation of plasmin within a fibrin clot from the plasminogen trapped within the

clot. Plasminogen-plasmin conversion, both within a clot and at its surface, is facilitated by the affinity of t-PA for fibrin, which results in a fibrin-dependent t-PA-induced plasminogen activation.

Plasminogen is a single-chain glycoprotein with a molecular weight of 5 92,000 which is synthesized by the liver and cleared from the circulation (via the liver) with a half-life of about 2.2 days. Human plasminogen comprises (i) a pre-activation peptide of about 67 to 76 amino-acids, (ii) five triple-loop disulfide bonded structures (named "kringles") of about 80 amino-acids, (iii) a catalytic serine proteinase unit of about 230 amino-acids, and (iv) some inter-domain 10 connecting sequences. Native plasminogen with NH₂-terminal glutamic acid (commonly named "Glu-plasminogen") is easily converted by limited digestion by plasmin of the Arg⁶⁸-Met⁶⁹, Lys⁷⁷-Lys⁷⁸, or Lys⁷⁸-Val⁷⁹ peptide bonds to modified forms commonly designated "Lys-plasminogen". Plasminogen is converted to plasmin by cleavage of the Arg⁵⁶¹-Val⁵⁶² peptide bond. The plasmin molecule is a 15 two-chain trypsin-like serine proteinase with an active site composed of His⁶⁰³, Asp⁶⁴⁶, and Ser⁷⁴¹. The kringles of plasminogen contain lysine binding sites that interact specifically with amino-acids such as lysine, 6-aminohexanoic acid and tranexamic acid. The lysine binding sites located in the kringle 1-3 region mediate the specific binding of plasminogen to fibrin and the kinetics of the interaction of 20 plasmin with α_2 -antiplasmin, and therefore play a crucial role in the regulation of physiological fibrinolysis.

Miniplasminogen is a derivative of plasminogen lacking the first four kringles which may be prepared by digestion of plasminogen with elastase and which is fully activatable to plasmin. It has a molecular weight of 38,000 and 25 contains over 100 amino-acids of the A chain including the fifth kringle structure.

Elevated pH conditions result in cleaving the Arg⁵³⁰-Lys⁵³¹ or Lys⁵³¹-Leu⁵³² bond of plasminogen and promoting disulfide bond rearrangement, thus producing microplasminogen, a derivative consisting of a 30 or 31 residue COOH-terminal peptide derived from the A chain bound through new disulfide 30 bonds to the intact B-chain of plasmin, as disclosed in U.S. Patent No. 4,774,087.

α_2 -antiplasmin is the main physiological plasmin inhibitor in human plasma which very rapidly inhibits plasmin, whereas plasmin formed in excess of α_2 -antiplasmin may be neutralized more slowly by macroglobulin and other serine proteinase inhibitors. α_2 -antiplasmin is a single-chain glycoprotein containing 464 amino acids which is present in plasma at a concentration of about 70 mg/l. During purification it is usually converted into a 452 amino-acid derivative by removal of 12 amino terminal amino-acids. α_2 -antiplasmin is synthesized by the liver and cleared from the circulation (via the liver) with a half-life of 2.6 days. Its reactive site is the Arg³⁷⁶-Met³⁷⁷ peptide bond. α_2 -antiplasmin is unique among serine proteinase inhibitors by having a COOH-terminal extension of 51 amino-acid residues which contains a secondary binding site that reacts with the lysine binding sites of plasminogen and plasmin. The native plasminogen-binding form of α_2 -antiplasmin becomes partly converted in the circulating blood to a non-plasminogen-binding, less reactive form, which lacks the 26 COOH-terminal residues. The Gln¹⁴-residue of α_2 -antiplasmin can crosslink to α -chains of fibrin by a process which requires Ca²⁺ and is catalyzed by activated coagulation factor XIII. α_2 -antiplasmin forms an inactive 1:1 stoichiometric complex with plasmin.

Plasmin or derivatives thereof (including mini- and microplasminogen), when infused in the vicinity of a clot in a dose sufficiently high to deplete α_2 -antiplasmin locally in an occluded blood vessel with stagnant flow, may have a sufficiently long half-life to be able to exert a local therapeutic effect. The administration of large amounts of plasmin is well tolerated, unlike the use of certain other proteolytic enzymes.

Thromboembolic disease, i.e. blockage of a blood vessel by a blood clot, affects many adults and can be a cause of death. Most spontaneously developing vascular obstructions are due to the formation of intravascular blood clots, known as thrombi. Small fragments of a clot (emboli) may detach from the body of the clot and travel through the circulatory system to lodge in distant organs and initiate further clot formation. Heart attack, stroke, renal and pulmonary infarcts

are well known consequences of thromboembolic phenomena. A blood clot is a gelled network of protein molecules within which are trapped circulating blood cells, platelets and plasma proteins. Fibrin is a major protein component of a clot which forms a relatively insoluble network. Proteolytic, particularly fibrinolytic 5 enzymes, have been used to dissolve vascular obstructions, since disruption of the fibrin matrix results in dissolution of the clot. Clots are formed when soluble fibrinogen, which is present in high concentrations in blood, is converted to insoluble fibrin by the action of thrombin. The probability of clot formation can be reduced by lowering the concentration of circulating fibrinogen, using 10 fibrinogenolytic enzymes.

Thromboembolic therapies have involved the administration of a plasminogen activator, e.g. either by direct intravenous injection, or by reinjection of a patient's plasma to which a plasminogen activator has been added *ex vivo*, or injection of plasma protein fractions previously mixed with streptokinase, or injection of 15 porcine plasmin stabilized with added lysine in conjunction with streptokinase.

Stroke is defined as a rapidly developing clinical sign of focal or global disturbance of cerebral function with symptoms lasting at least 24 hours. Stroke is typically caused by blockage or occlusion of blood vessels to the brain or within the brain. With complete occlusion, arrest of cerebral circulation causes cessation 20 of neuronal electrical activity within seconds. Within a few minutes after deterioration, depletion of high energy phosphates, membrane ion pump failure, efflux of cellular potassium, influx of sodium chloride and water, and membrane depolarization occur. If the occlusion persists for more than five to ten minutes, irreversible damage results. With incomplete ischemia, however, the outcome is 25 difficult to evaluate and depends largely on residual perfusion and the availability of oxygen. After a thrombotic occlusion of a cerebral vessel, ischemia is rarely total. Some residual perfusion usually persists in the ischemic area, depending on collateral blood flow and local perfusion pressure.

Cerebral blood flow can compensate for drops in mean arterial blood 30 pressure from 90 to 60 mm Hg by auto-regulation. This phenomenon involves

dilatation of downstream resistant vessels. Below 60 mm Hg, vasodilatation is inadequate and the cerebral blood flow falls. The brain however has perfusion reserves that can compensate for the fall in cerebral blood fall. When distal blood pressure falls below about 30 mm Hg, both compensatory mechanisms (auto-
5 regulation and perfusion reserve) are inadequate to prevent failure of oxygen delivery. As flow drops below the ischemic threshold, symptoms of tissue hypoxia develop. Severe ischemia may be lethal. Moderate ischemia results in a tissue area that can be saved called penumbra. In the neurological context, penumbra refers to a zone of brain tissue with moderate ischemia and paralyzed neuronal
10 function, which is reversible with restoration of adequate perfusion. The penumbra forms a zone of collaterally perfused tissue surrounding a core of severe ischemia in which an infarct has developed. When a clot is degraded and the blood flow to the penumbra is restored, the phenomenon of reperfusion injury can occur.

15 Although an ischemic event can occur anywhere in the vascular system, the carotid artery bifurcation and the origin of the internal carotid artery are the most frequent sites for thrombotic occlusions of cerebral blood vessels, which result in cerebral ischemia. The symptoms of reduced blood flow due to stenosis or thrombosis are similar to those caused by middle cerebral artery disease. Flow
20 through the ophthalmic artery is often affected sufficiently to produce transient monocular blindness. Severe bilateral internal carotid artery stenosis may result in cerebral hemispheric hypoperfusion. This manifests with acute headache ipsilateral to the acutely ischemic hemisphere. Occlusions or decrease of the blood flow with resulting ischemia of one anterior cerebral artery distal to the
25 anterior communicating artery produces motor and cortical sensory symptoms in the contralateral leg and, less often, proximal arm. Other manifestations of occlusions or underperfusion of the anterior cerebral artery include urinary incontinence due to damage to the parasagittal frontal lobe. Language disturbances manifested by decreased spontaneous speech may accompany
30 generalized depression of psychomotor activity.

Most ischemic strokes involve portions or all of the territory of the middle cerebral artery, with emboli from the heart or extracranial carotid arteries accounting for most cases. Emboli may occlude the main stem of the middle cerebral artery, but more frequently produce distal occlusion of either the superior 5 or the inferior branch. Occlusions of the superior branch cause weakness and sensory loss that are greatest in the face and arm. Occlusions of the posterior cerebral artery distal to its penetrating branches cause complete contra-lateral loss of vision. Difficulty in reading (dyslexia) and performing calculations (dyscalculia) may follow ischemia of the dominant posterior cerebral artery. 10 Proximal occlusion of the posterior cerebral artery causes ischemia of the branches penetrating to calamic and limbic structures, resulting in disturbances that may chronically change to intractable pain of the defective site (thalamic pain).

A significant event in cerebral ischemia is known as the transient ischemic 15 attack ("TIA"), defined as a neurologic deficit with a duration of less than 24 hours. TIA is an important sign of an ischemic development that may lead to cerebral infarction. Its etiology involves hemodynamic events and thromboembolic mechanisms. Because TIA often resolves within one hour, a longer deficit is often classified as presumptive stroke and is, accordingly, 20 associated with permanent brain injury. Therefore, computed tomographic brain scans are used to search for cerebral infarction in areas affected by TIA lasting longer than two hours. Thus, the relevant clinical distinction between TIA and stroke is whether ischemia has caused brain damage, which is typically classified as infarction or ischemic necrosis. Subjects with deteriorating clinical signs might 25 have stroke in evolution (progressive stroke).

Many other diseases are caused by or associated with ischemia. For instance, vertebrobasilar ischemia results from occlusion of the vertebral artery which causes lateral medullary syndrome with symptoms including vertigo, nausea, ipsilateral ataxia and Herner's syndrome. Vertebrobasilar ischemia often 30 produces multifocal lesions scattered on both sides of the brain stem along a

considerable length. A basilar artery occlusion produces massive deficits, including paralysis of the limbs and of most bulbar muscles, leaving the subject only able to communicate by moving the eyes or eyelids and producing an initial reduction in arousal followed by blindness and amnesia.

5 Venous occlusion can cause massive damage and death. The primary mechanism of brain damage is then a reduction in capillary blood flow because of increased outflow resistance from the blocked veins. Back transmission of high pressure into the capillary bed usually results in early brain swelling from oedema and hemorrhagic infarction in subcortical white matter. The most dangerous form
10 of venous disease arises when the superior sagittal sinus is occluded. Venous occlusion occurs in association with coagulation disorders, often in the purpural period or in subjects with disseminated cancers.

15 Brief diffuse cerebral ischemia can cause syncope without any permanent sequel. Prolonged diffuse ischemia in other organs has devastating consequences. Common causes are cardiopulmonary failure, including infarction, aortic dissection and global hypoxia or carbon monoxide poisoning. Clinically, a diffuse hypoxia/ischemia results in unconsciousness and coma, often followed by a chronic vegetative state. If the subject does not regain consciousness within a few days, chances for the return of independent brain functions becomes very
20 poor.

25 Hyperviscosity syndrome is another disease related to blood flow and ischemia. Subjects with hyperviscosity syndrome can present either with focal neurologic dysfunction or with diffuse or multifocus signs or symptoms including headache, visual disturbances, cognitive impairments or seizures.

Ischemic stroke due to thrombotic closure of a cerebral artery is amenable to therapy with antithrombotic and thrombolytic agents. The use of t-PA within three hours of symptom onset is associated with a better neurologic outcome, but a significant percentage of treated patients experience acute hemorrhage in the brain. Thus, the development of safer and more effective treatments is needed.

US-A-5,288,489 discloses a method of dissolving an intravascular thrombus in a human patient, or reducing the risk of thrombus formation in a patient (such as diabetics and pregnant women), comprising administering parenterally to the patient a therapeutically effective amount of human or 5 mammalian plasmin or mini-plasmin or micro-plasmin in a fibrinolytic or fibrinogenolytic active form, the said active form being obtained either by exposure to an insolubilized, entrapped, encapsulated or immobilized plasminogen activator or by inhibiting the autolytic activity by means of certain hydrophobic ions. This method is disclosed in the context of heart attack, stroke, 10 renal and pulmonary infarctions, thrombophlebitis, and so on. EP-A-631,786 discloses administration to a subject of a protein having the effect of lys- plasminogen for the treatment of ischemia, infarction, brain edema and reperfusion injury that follows ischemic events. WO 00/18436 discloses the use 15 of plasmin, mini-plasmin or micro-plasmin in a therapeutic composition for the treatment of focal cerebral ischemic infarction (ischemic stroke).

In the population over 60 years of age, the prevalence of intermittent claudication or chronic peripheral arterial occlusive disease (PAOD) being the result of atherosclerotic and thrombotic processes, is between 1 and 8%. Over the course of their disease, about 20% of the patients with intermittent 20 claudication will progress to critical leg ischemia (acute PAOD) endangering the viability of the lower extremity, 10% will undergo invasive/surgical procedures for progressive symptoms, and 5% require amputation of the limb. Blood flow can be restored through operative bypass surgery, vascular repair surgery or pharmacological dissolution of the blood clot. Intra-arterial thrombolysis is 25 expected to provide a significant reduction in surgical procedures, without increased risk of amputation or death. Urokinase is currently the most widely used agent for intra-arterial thrombolysis.)

Plasminogen can be obtained from human plasma fractions by affinity chromatography on lysine-Sepharose, however with yields of no more than 0.25 30 g/l. With the general reluctance to use plasma fractionation derivatives,

alternative approaches such as production via recombinant DNA technology are preferred. For the production of a large and complex molecule such as plasminogen or plasmin, however, an effective expression system is required. Indeed recombinant intact plasminogen cannot readily be expressed in 5 activatable form in common eukaryotic expression systems, due to the nearly ubiquitous presence of intracellular plasminogen activators within such cell types, resulting in degradation of human plasminogen in the conditioned cell culture media. According to J. Wang et al. in *Protein Science* (1995) 4:1758-1767, a baculovirus/insect cell expression system has enabled expression of 10 microplasminogen at low levels of 3 to 12 mg/l. Such a yield is obviously too low for the production of large quantities of the purified active substance. We are not aware of any data relating to the expression of miniplasminogen. Therefore, there is a need in the art for an expression system making it possible to produce large amounts of plasminogen and derivatives thereof, including mini- and 15 microplasminogen, which will be useful in the treatment of ischemic and thrombotic disorders and associated diseases such as listed hereinabove.

SUMMARY OF THE INVENTION

The present invention relates to the use of certain yeast, e.g. *Pichia pastoris*, for the high yield production of recombinant mammalian plasminogen 20 and derivatives thereof (including, but not limited to, miniplasminogen and to microplasminogen) and to the production of recombinant mammalian plasmin and derivatives thereof in sufficient amounts, purity and stability to be clinically applicable for the treatment of mammal, specifically humans and horses. The therapeutic efficacy of the recombinant human microplasmin obtained according 25 to this production method was illustrated in animal models of ischemic stroke, acute myocardial infarction and extracorporeal arteriovenous circulation thrombosis models.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 is a schematic representation of the amino-acid structure of plasminogen, wherein black bars indicate disulphide bonds; Pli is a plasmic cleavage site for conversion of Glu-plasminogen to Lys-plasminogen; UK is the cleavage site for plasminogen activators, yielding plasmin; μ PIg and mPIg 5 respectively indicate the origin of microplasminogen and miniplasminogen used in this invention.

Figure 2 shows individual data of angiographic examinations in dogs with a copper coil induced thrombosis after treatment with the recombinant microplasminogen of this invention.

10 Figure 3 shows the nucleotide sequence (SEQ. ID. No. 3) and amino-acid sequence (SEQ. ID. No. 4) of human microplasminogen.

Figure 4 shows the nucleotide sequence (SEQ. ID. No. 5) and amino-acid sequence (SEQ. ID. No. 6) of human miniplasminogen.

DEFINITIONS

15 The term "catalytic domain of plasminogen", as used herein, refers to a serine protease unit of about 230 amino-acids that, after activation by a plasminogen activator, digest fibrin to soluble degradation products.

20 The term "mutants", as used herein, refers to a catalytically active protein sequence in which one or more amino-acids are substituted, deleted or mutated, and wherein the level of similarity with the wild-type protein is at least 80%, preferably at least 85% and more preferably at least 90%.

25 The term "hybrids", as used herein, refers to the protein of a mammal wherein at least one sequence of one or more amino-acids is replaced by a sequence, preferably the corresponding sequence, from the corresponding protein of another mammal. With respect to plasminogen, the sequence replacement may be either in the catalytic domain or in any of the five kringle domains.

DETAILED DESCRIPTION OF THE INVENTION

A first object of the invention is a yeast expression vector comprising a mammalian nucleotide sequence operably linked to a promoter, wherein the said mammalian nucleotide sequence codes for the catalytic domain of plasminogen and further optionally codes for one or more kringle domains of plasminogen, or mutants or hybrids thereof. In preferred embodiments of the said yeast expression vector, the mammalian nucleotide sequence codes for plasminogen, microplasminogen and miniplasminogen respectively. In a more specific embodiment, the nucleotide sequence is a human nucleotide sequence, such as SEQ ID No.1 (plasminogen), SEQ. ID. No. 3 (miniplasminogen) or SEQ. ID. No.5 (microplasminogen). In a preferred embodiment, the promoter is an inducible promoter. In another preferred embodiment, the yeast expression vector is able to stably integrate in the yeast genome, e.g. by homologous recombination. In preferred embodiments, the nucleotide sequence is fused to a secretion signal, i.e. a peptide that targets protein to the cell membrane where the signal peptide is cleaved and the protein released in the medium, for example α -factor, PHO or AGA-2.

Preferably, the yeast expression vector of the invention is for expression in a yeast selected from the group consisting of methylotrophic yeasts represented by the genera *Hansenula*, *Pichia*, *Candida* and *Torulopsis*.

A second object of the invention is a yeast cell comprising a mammalian nucleotide sequence coding for the catalytic domain of plasminogen and further optionally coding for one or more kringle domains of plasminogen, or mutants or hybrids thereof. In preferred embodiments of the said yeast cell, the mammalian nucleotide sequence codes for plasminogen, microplasminogen and miniplasminogen respectively, i.e. the yeast cell comprises the nucleotide sequence SEQ.ID.No. 1 or the nucleotide sequence SEQ.ID.No. 3 or the nucleotide sequence SEQ.ID.No. 5.

The invention also relates to a yeast cell transfected with a vector such as disclosed hereinabove. Preferably, the said vector is integrated in the genome.

The yeast cell of the invention preferably belongs to the group of methylotrophic yeasts. More particularly, the said yeast may be selected from the genera consisting of, *Pichia*, *Hansenula*, , *Candida*, and *Torulopsis*. In another embodiment, the yeast cell of the invention belongs to the *Pichia pastoris* 5 species.

An exemplary yeast cell belongs to the cell line deposited with the Belgian Coordinated Collections of Micro-organisms under accession number MUCL43676. A most significant advantage of the yeast cells of the invention is their ability to express human microplasminogen at a level of at least about 100 10 mg/litre, i.e. at a much higher level than was known in the art. Another advantage is their ability to express human miniplasminogen at a level of at least about 3 mg/litre.

Another object of the invention is a method of expressing a mammalian protein comprising the catalytic domain of plasminogen and further optionally 15 comprising one or more kringle domains of plasminogen, or mutants or hybrids thereof, in a yeast cell such as defined hereinabove, using recombinant technology procedures well known in the art which are detailed in the appended examples. Specific embodiments of the said method relate to mammalian proteins having respectively the amino-acid sequence SEQ.ID. No. 2 (human 20 plasminogen), the amino-acid sequence SEQ.ID. No. 4 (human microplasminogen) or the amino-acid sequence SEQ.ID. No. 6 (human miniplasminogen). In a preferred embodiment, the method further comprises the step of activating the expressed mammalian protein by means of a plasminogen activator which may be staphylokinase or a variant thereof. In another preferred 25 embodiment, the method further comprises the step of stabilizing the expressed and activated mammalian protein by means of a stabilizing agent. The said stabilizing agent may comprise either an amino-acid selected from the group consisting of lysine, 6-amino hexanoic acid and tranexamic acid or a stabilizing medium. The latter may suitably be an acid solution or an acid buffer such as a 30 citrate buffer with a pH of about 3.1. Still preferred is a method further comprising

the step of drying the expressed, activated and stabilized mammalian protein, e.g. by means of lyophilization.

The present invention also relates to a recombinant mammalian protein obtained by a method such as described hereinabove or expressed in a yeast cell 5 such as defined hereinabove. This protein is obtainable in large amounts and in a high purity level, thus meeting the necessary requirements for being used as an active ingredient in pharmaceutical compositions for the treatment of the various ischemic diseases listed in the above section "background of the invention".

The present invention will be demonstrated in more detail in the following 10 examples, which are however not intended to be limiting the scope of the invention, the latter being only defined by the appended claims.

EXAMPLE 1 - vector construction for expression of human microplasminogen and human miniplasminogen in *Pichia pastoris*

The pPICZ α A secretion vector purchased from Invitrogen Corporation 15 (Carlsbad, California) was used to direct expression and secretion of recombinant human microplasminogen in *Pichia pastoris*. Relevant features of this vector are:

- a 942 bp fragment containing the alcohol oxidase 1 (AOX1) promoter that allows methanol-inducible, high level expression in *Pichia* and targeted plasmid integration to the AOX1 chromosomal locus,
- the native transcription termination and polyadenylation signal from the AOX1 gene;
- an expression cassette conferring zeocin resistance to *Escherichia coli* and *Pichia pastoris*;
- a ColE1 origin of replication for propagation and maintenance of the plasmid in *E. coli*, and
- unique restriction sites (Sacl, Pmel, BstXI) that permit linearization of the vector at the AOX1 locus for efficient integration into the *Pichia* genome.

In addition to the above features, this vector contains the secretion signal of the *Saccharomyces cerevisiae* α -factor prepropeptide, allowing expression of heterologous proteins as secreted proteins in the medium. The processing of the α factor mating signal sequence in pPICZ α occurs in two steps:

- 5 1. the preliminary cleavage of the signal sequence by the *KEX2* gene product, with the Kex2 cleavage occurring between arginine and glutamine in the sequence Glu-Lys-Arg * Glu-Ala-Glu-Ala, wherein * is the site of cleavage.
2. the Glu-Ala repeats are further cleaved by the *STE13* gene product.

10 However, the Glu-Ala repeats are not always necessary for cleavage by Kex2, depending on the amino acid following the Glu-Lys-Arg sequence. In some cases where Ste13 cleavage is not efficient, the Glu-Ala repeats are left on the NH₂-terminus of the expressed protein of interest.

A Xhol recognition sequence is present at the COOH-terminus of the α factor secretion signal, immediately upstream of the Lys-Arg Kex2 cleavage site. This Xhol restriction site may be used to clone the gene of interest flush with the Kex2 cleavage site by using a PCR cloning approach and an appropriate forward primer to rebuild the sequence from the Xhol site to the arginine codon. The recombinant protein of interest will then be expressed with a native NH₂-terminus.

20 Engineered immediately downstream of the α factor signal sequence in the pPICZ α A vector is a multi-cloning site with recognition sequences for the enzymes EcoRI, Sfil, KpnI, Xhol, SacII and XbaI to facilitate the cloning of foreign genes.

Expression vector construction for microplasminogen

25 The vector Fmyc- μ Pli disclosed by Lasters et al. in *Eur. J. Biochem.* (1997) 244:946 was used to isolate, by amplification ("PCR-rescue") with the Advantage cDNA polymerase mix available from Clontech (Palo Alto, California), the region encoding the human microplasminogen protein. After a DNA template

denaturation step of 3 minutes at 94°C, 30 temperature cycles were performed (30 seconds at 94°C, 30 seconds at 50°C, 30 seconds at 72°C), followed by a 2 minutes final elongation step at 72°C. The following oligonucleotide primers LY-MPG1 (sense) and LY-MPG2 (antisense) were used in this reaction:

5 LY-MPLG1: 5' GGGGTATCT CTC GAG AAA AGA GCC CCT TCA TTT GAT TG
(SEQ.ID. No. 7)

LY-MPLG2: 5' GTTTTTGT TCT AGA TTA ATT ATT TCT CAT CAC TCC CTC
(SEQ.ID. No. 8)

The LY-MPLG1 primer had an annealing region corresponding to residues 10 543-548 of plasminogen (Ala-Pro-Ser-Phe-Asp-Cys) preceded by a non-annealing extension which included the last four residues of the α factor mating signal (Leu-Glu-Lys Arg). In this extension, the Leu-Glu codons determine the Xho I restriction site (underlined) allowing the cloning of the gene of interest flush with the Kex2 cleavage site. The LY-MPLG2 primer had an annealing region 15 corresponding to the last seven residues of plasminogen, followed by a TAA stop-codon and a non-annealing region comprising a XbaI recognition sequence.

The amplified fragment having the expected size (~ 780 bp) was digested with Xhol and XbaI, and directionally cloned into the vector pPICZ α A. The recipient vector-fragment was prepared by Xhol and XbaI restriction, and purified 20 from agarose gel using the Qiaquick gel extraction kit (Qiagen GmbH, Germany). The E.coli strain TG1 (DSMZ collection #1208, Germany) was transformed with the ligation mixture, and zeocin resistant clones were selected. Based on restriction analysis, a plasmid clone containing an insert of the expected size was retained for further characterization. Sequence determination of the vector 25 pPICZ α -MPLG1 (clone #5) confirmed the precise insertion of the microplasminogen coding region fused to the α factor mating signal, as well as the absence of unwanted mutations in the coding region. The primers 5'AOX and 3'AOX were provided in the EasySelect *Pichia* expression kit from Invitrogen, Carlsbad, California.

The determined nucleotide sequence and the deduced amino-acid sequence of human microplasminogen used are represented in SEQ.ID No. 3 and SEQ.ID No. 4, respectively. Compared to the sequence previously determined by Forsgren et al. in *FEBS Lett.* (1987) 213: 254, the nucleotide 5 sequence differs in 10 positions. However, the amino acid sequence was identical.

Expression vector construction for miniplasminogen

A pPICZ α -derived secretion vector was constructed as follows for miniplasminogen expression, making use of the hereinabove described 10 pPICZ α -MPLG1 vector.

The vector FdTet-SN-miniPIg disclosed by Lasters et al. (cited *supra*) was used to isolate by amplification ("PCR-rescue") a 500 bp DNA fragment encoding kringle five and part of the catalytic domain of the miniplasminogen protein. After a DNA template denaturation step of 3 minutes at 94°C, 30 temperature cycles 15 were performed (10 seconds at 94°C, 10 seconds at 50°C, 15 seconds at 72°C), followed by a 2 minutes final elongation step at 72°C. The following oligonucleotide primers LY-MINPLG1 (sense) and LY-MINPLG2 (antisense) were used in this reaction:

LY-MINPLG1: 5' GGGGTATCT CTC GAG AAA AGA GCA CCT CCG CCT GTT
20 GTC CTG CTT CC (SEQ.ID. No. 9)

LY-MINPLG2: 5' GCA GTG GGC TGC AGT CAA CAC CCA CTC (SEQ.ID. No. 10)

The LY-MINPLG1 primer has an annealing region corresponding to residues 444-452 of plasminogen (Ala-Pro-Pro-Pro-Val-Val-Leu-Leu-Pro) 25 preceded by a non-annealing extension which included the last four residues of the factor mating signal (Leu-Glu-Lys-Arg). In this extension, the Leu-Glu codons determine the Xho I restriction site allowing the cloning of the gene of interest flush with the Kex2 cleavage site.

The LY-MINPLG2 primer has an annealing region corresponding to the residues 596-604 of plasminogen (Glu-Trp-Val-Leu-Thr-Ala-Ala-His-Cys). This annealing region of the catalytic domain, also present in the microplasminogen expression vector, comprises a unique Pst I recognition sequence (underlined).

5 The amplified fragment having the expected size was digested with Xhol and PstI, and directionally cloned into the vector pPICZ α -MPLG1 described above (microplasmin expression vector). The recipient vector-fragment was prepared by Xhol and PstI restriction, and purified from agarose gel using the Qiaquick gel extraction kit (Qiagen GmbH, Germany). The *E. coli* strain TG1
10 (DSMZ collection #1208, Germany) was transformed with the ligation mixture, and zeocin resistant clones were selected. Based on restriction analysis, a plasmid clone containing an insert of the expected size was retained for further characterization. Sequence determination of the vector pPICZ α -KMPLG1 (clone #3) confirmed the precise insertion of the amplified fragment fused to the α -factor
15 mating signal, as well as the absence of unwanted mutations in the cloned region (the primers LY-MINPLG1 and LY-MINPLG2 were used).

EXAMPLE 2 - high level expression and purification of recombinant human microplasminogen: quantitative activation and stabilization of microplasmin

10 μ g of the vector pPICZ α -MPLG1 was digested with PmeI, which
20 linearizes the vector in the 5' AOX1 region. The DNA was concentrated to about 0.33 μ g/ μ l by precipitation, and 5 μ l was used to transform competent *Pichia pastoris* X33 cells prepared according to the manual provided in the EasySelect
Pichia expression kit.

The selection of a high-expression strain was performed as follows. Zeocin
25 resistant transformants were selected on YPDSZ plates (1% yeast extract, 2% peptone, 2% glucose, 1M sorbitol, 2% agar, 100 μ g/ml zeocin). Thirty-four single colonies were inoculated in 10 ml BMYZ-glycerol medium (1% yeast extract, 2% peptone, 1% glycerol, 100 mM potassium phosphate, pH 6.0, 1.34% yeast nitrogen base, 4x10⁻⁵ % biotin, 100 μ g/ml zeocin) in 50 ml Falcon tubes and

cultured for 16 hours at 30°C. The cells were pelleted and re-suspended in 2 ml of BMYZ-methanol medium (same as BMYZ-glycerol but with 0.5% methanol instead of glycerol) to induce expression from the AOX1 promoter, and cultured for 40 hours. 4 pulses of 0.5 % methanol were regularly supplied to the cultures
5 over this period. At the end of the induction culture, the presence of microplasminogen in the culture supernatant was estimated as described by Lijnen et al. in *Eur. J. Biochem.* (1981) 120:149. Briefly, the microplasminogen in pure or 10-fold diluted supernatants were incubated with urokinase for 30 minutes to activate microplasminogen in microplasmin. The generated
10 microplasmin activity, as determined by its amidolytic activity measured with the chromogenic substrate S2403 (available from Chromogenix, Antwerp, Belgium) at different times, was compared to the activity of known amounts of purified plasmin or microplasmin preparations. The clone X33-MPLG1 #5, showing the highest microplasmin activity after urokinase activation, was selected for
15 subsequent large scale production.

Fermentation of X33-MPLG1#5 at a 50 litre scale was carried out in four steps as follows. 2 l flask cell cultures were performed for 23 hours at 30°C in 400 ml YSG+ (yeast extract 6 g/l, soya peptone 5 g/l, glycerol 20 g/l) using an inoculum of 0.7 ml of cell bank (glycerol OOC17) and 270 rpm agitation, yielding
20 (at the end of the pre-culture step) an OD600 of 15. Fermentation was then performed in a MRP80 fermentation device in 30 l basal medium (26.7 ml/l H₃PO₄ 85%; 1.05 g/l CaSO₄.2H₂O, 18.2 g/l K₂SO₄, 14.9 g/l MgSO₄.7H₂O, 4.13 g/l KOH, 40 g/l glycerol 100% and 4.76 ml/l PTM1 salt solution [comprising 6 g/l CuSO₄.5H₂O, 0.08 g/l NaI, 3.36 g/l MnSO₄.H₂O, 0.2 g/l NaMoO₄.2H₂O, 0.02 g/l
25 Boric acid, 0.82 g/l CoCl₂.6H₂O, 20 g/l ZnCl₂, 65 g/l FeSO₄.7H₂O, 0.2 g/l d-biotin and 5 ml/l HSSO₄]), using 600 ml inoculum at 30°C with an air flow of 50 l/min at atmospheric pressure, dissolved oxygen (DO) >20% and 200-500 rpm agitation, pH being maintained at 5.8 with 12.5% ammonia. At 24 hours and OD 600 of 50 (end of batch step), glycerol depletion was evidenced by a rapid
30 increase of dissolved oxygen. Glycerol feeding (632 g/l glycerol 100% and 12 ml/l

PTM1) increased the OD 600 up to 258 in 24 hours. Methanol feeding was then carried out with an increasing flow of up to 250 ml/h within 6 hours, which was maintained for 66 hours using 988 ml/l methanol and 12 ml/l PTM1 to reach an OD 600 of 352 at the end of culture. Fermentation of X33-MPLG1#5 at a 350 litre

5 scale provided proportionally similar results.

The harvest was then purified in a three-steps process comprising cation exchange expanded bed chromatography, hydrophobic chromatography and affinity chromatography as follows:

a) Cation exchange expanded bed chromatography

10 Cation exchange expanded bed adsorption chromatography was conducted with Streamline SP (available from Pharmacia Biotechnology, Cat. No. 17-0993-01/02) packed in a Streamline 200 column (Pharmacia Biotechnology Cat No. 18-1100-22) with a bed volume of 5,120 cm³, expanded and equilibrated by applying an upward flow of 1 M NaCl, 25 mM sodium acetate (CH₃COONa.3

15 H₂O), buffer, pH 6.0, for two column volumes followed by column volumes of 25 mM sodium acetate buffer, pH 6.0. The fermentation broth was on line diluted (7x) with water and passed upwards through the expanded bed at a flow rate of 1000 ml/min. Loosely bound material was washed out with the upward flow of 25 mM sodium acetate buffer pH 6.0. The column adaptor was then lowered to the

20 surface of the sedimented bed at a height of 16.3 cm. Flow was reversed and the captured proteins eluted with 2 column volumes of 0.5 M NaCl, 25 mM sodium acetate buffer, pH 6.0. Solid ammonium sulfate was added to the eluted Streamline fraction to reach 30 % saturation (164 g ammonium sulfate per liter of eluted Streamline fraction) and the mixture was gently stirred at 4 - 8°C for 1

25 hour.

b) Hydrophobic chromatography

Hydrophobic chromatography was conducted with Hexyl TSK 650C (available from Toso-Haas Cat. No. 19027) packed in a Vantage 180/500 column (available from Millipore, Cat. No. 87018001) with a packed volume of 2,700 cm³

at 4-8°C. The eluted streamline fraction was loaded on the column at a flow rate of 38 l/hour. The column was then washed with 1.5 column volumes of 25 mM sodium acetate buffer, pH 6.0, containing 164 g/l ammonium sulfate and eluted from the column with 7 column volumes of 25 mM sodium acetate buffer, pH 6.0.

5 c) Affinity chromatography

Affinity chromatography was conducted with Blue Sepharose 6 Fast Flow (available from Pharmacia Biotechnology, Cat. No. 17-0948-02/03) packed in a Vantage 130/500 column (available from Millipore, Cat. No. 87013001) with a packed volume of 3,186 cm³ at 4-8°C. The eluted fraction was loaded on the 10 column at a flow rate of 20 l/hour, and washed with one column volume of 25 mM disodium hydrogenophosphate (Na₂HPO₄.12 H₂O) buffer, pH 7.0. The microplasminogen protein fraction was eluted from the column with 5 column 15 volumes 0.5 M NaCl, 25 mM di-sodium hydrogenophosphate buffer, pH 7.0 and kept frozen at -20°C. The purity of the material was above 98% as demonstrated by SDS gel electrophoresis.

Quantitative activation to and stabilization of microplasmin

a) Quantitative activation

The activation of microplasminogen to microplasmin was performed at 23°C for 30 minutes at a molar ratio of 0.5 % of a staphylokinase variant SY162 in 0.5 M 20 NaCl, 25 mM di-sodium hydrogenophosphate (Na₂HPO₄.12 H₂O) buffer, pH 7.0. SY162 is a staphylokinase variant with reduced immunogenicity comprising 12 amino-acid substitutions (K35A, E65Q, K74R, E80A, D82A, T90A, E99D, T101S, E108A, K109A, K130T and K135R) as compared to wild-type, as described by WO 99/40198. Solid ammonium sulfate was added to microplasmin at a final 25 concentration of 1 M (132 g/l) and the mixture stirred at 4 - 8°C for 15 minutes.

b) Hydrophobic chromatography

Hydrophobic chromatography was conducted with Phenyl Sepharose 6 Fast Flow (available from Pharmacia Biotechnology, Cat. No. 17-0965-03/05) packed

in a BPG 100/500 column (available from Pharmacia Biotechnology, Cat. No. 18-1103-01) having a packed volume of 1,738 cm³, equilibrated with 4 column volumes of 25 mM Na₂HPO₄.12 H₂O buffer, pH 7.0, containing 0.1 M tranexamic acid (available from Bournonville Pharma, Braine-L'Alleud, Belgium) and 1 M (NH₄)₂SO₄, pH 7.0, at 4-8°C. The activated product was loaded on the column at a linear flow rate of 18 l/hour and washed with 4.5 column volumes of 25 mM Na₂HPO₄.12 H₂O buffer, pH 7.0, containing 0.1 M tranexamic acid and 1 M (NH₄)₂SO₄. Microplasmin was eluted from the column at a linear flow rate of 6 l/hour with 5 column volumes of 25 mM Na₂HPO₄.12 H₂O buffer, pH 7.0, containing 0.1 M tranexamic acid and 0.7 M (NH₄)₂SO₄ and equilibrated with phosphate buffered saline containing 0.1 M tranexamic acid. Staphylokinase variant SY162 was eluted from the column with 25 mM Na₂HPO₄.12H₂O buffer, pH 7.0 containing 0.1 M tranexamic acid. This procedure removed above 99% of staphylokinase from the microplasmin peak as demonstrated with a specific 15 ELISA assay.

c) Concentration and diafiltration by tangential ultrafiltration

Ultrafiltration was conducted with 2 Pellicon 2 Biomax membranes (5 kDa, 2.5 µm, available from Millipore, Bedford, Massachussets, Cat. n° P2B005A25) at 2-8°C. The membranes were mounted in a Pellicon 2 Process Holder connected 20 to a Microgon pump Cart System (available from Microgon, Laguna Hills, California). The membranes were washed with purified water and membrane integrity tested before operation. The sanitization was performed by continuous recirculation with 0.5 M NaOH for 60 minutes and with 0.1 M NaOH during 60 minutes. The membranes were then rinsed with 5 mM citric acid, pH 3.1, until the 25 permeate reached a pH of 3.1. The pH of the Phenyl Sepharose eluate was adjusted to 3.1 and the protein was concentrated to 4 mg/ml by ultrafiltration. Diafiltration was performed for 60 to 90 minutes against 5 volumes of 5 mM citric acid, pH 3.1. Yields (expressed in grams) of three runs performed on a 50 litre 30 fermentation apparatus are summarized in the following Table 1 (ND : not determined).

Table 1

	Run 1	Run 2	Run 3
Fermentor	220	240	ND
Streamline	50	79	130
Hexyl	36	37	ND
Blue	25	28	30
Phenyl	17	20	26
Diafiltration	-	-	22

d) Sterile filtration (0.2 µm)

5 Mannitol was added at 2-8°C to a concentration of 1.5 g/g of protein and sterile filtration performed at 23°C on a Millipak 100 filter (size 500 cm²) (available from Millipore, Cat. No. MPGL10CA3) and rinsed with about 500 ml of 5 mM citric acid, pH 3.1, with a peristaltic pump at a flow rate of 500 ml/minute. The filtrate was collected in a sterile and pyrogen free bag and stored at -20°C.

EXAMPLE 3 - expression of recombinant human miniplasminogen

10 About 15 µg of the vector pPICZ α -KMPLG1 was digested in a 20 µl reaction with Pmel, which linearizes the vector in the 5' AOX1 region. The linear DNA (3 µg) was used to transform competent *Pichia pastoris* X33 cells prepared according to the manual provided in the EasySelect *Pichia* Expression kit.

15 The selection of high-expression strain was performed essentially as follows. Zeocin resistant transformants were selected on YPDSZ plates (as defined in example 2). Fifty isolated colonies were inoculated in 15 ml BMYZ-glycerol medium (as defined in example 2) in 50 ml Falcon tubes and cultured for

16 hours at 30°C. The cells were pelleted and re-suspended in 1.5 ml of BMYZ-methanol medium (as defined in example 2) to induce expression from the AOX1 promoter, and cultured for 40 hours. 3 or 4 pulses of 0.5 % methanol were regularly supplied to the cultures over this period. At the end of the induction 5 culture, the presence of miniplasminogen in the culture supernatant was estimated as described by Lijnen et al. (cited *supra*). Briefly, the miniplasminogen in 10-fold diluted supernatants were incubated with streptokinase for 10 minutes to form an active complex. The generated miniplasmin activity, as determined with the chromogenic substrate S2403 (see example 2) at different times, was 10 compared to the activity of known amounts of a purified plasminogen preparation. In these conditions, all tested clones were producing miniplasminogen with yields varying between 3 and 15 mg/l. The two clones X33-KMPLG1 #6 and X33-KMPLG1 #25, showing the highest miniplasmin activity, were selected for subsequent large scale production.

15 **EXAMPLE 4 - murine cerebral ischemic infarction model (general procedure)**

Experiments were conducted according to the guiding principles of the American Physiological Society and the International Committee on Thrombosis and Haemostasis as disclosed by Giles in *Thromb. Haemost.* (1987) 58:1078.

Focal cerebral ischemia was produced by persistent occlusion of the 20 middle cerebral artery (hereinafter MCA) according to Welsh et al. in *J. Neurochem.* (1987) 49:846. Briefly, mice of either sex, weighing 20 to 30 g, were anesthetized by intraperitoneal injection of 75 mg/ml ketamine (available from Apharmo, Arnhem, The Netherlands) and 5 mg/ml xylazine (available from Bayer, Leverkusen, Germany). Alternatively, in order to ensure that these drugs did not 25 affect cerebral infarct size, anesthesia was performed with inhalation of 2% isoflurane in oxygen. 1 mg/kg atropine (available from Federa, Brussels, Belgium) was administered intramuscularly, and rectal temperature was maintained at 37°C by keeping the animals on a heating pad. A "U" shape incision was made between the left ear and left eye. The top and backside segments of the temporal 30 muscle were transected and the skull was exposed by retraction of the temporal

muscle. A 1 mm diameter opening was made in the region over the MCA with a hand-held drill, with saline superfusion to prevent heat injury. The meninges were removed with a forceps and the MCA was occluded by ligation with a 10-0 nylon thread (available from Ethylon, Neuilly, France) and trans-sected distally to the 5 ligation point. Finally the temporal muscle and skin were sutured back in place. The recombinant microplasmin produced in example 2 was then given intravenously as a bolus, 15 minutes after ligation of the MCA unless otherwise indicated. The animals were allowed to recover. After 24 hours, the animals were sacrificed with 500 mg/kg Nembutal (available from Abbott Laboratories, North 10 Chicago, Illinois) and decapitated. The brain was removed and placed in a matrix for sectioning in 1 mm segments. The sections were immersed in 2% 2,3,5-triphenyltetrazolium chloride in saline, incubated for 30 minutes at 37°C, and placed in 4% formalin in phosphate buffered saline. With this procedure, the necrotic infarct area remains unstained and is clearly distinguishable from stained 15 viable tissue. The sections were photographed and subjected to planimetry. The volume of the focal cerebral ischemic injury was defined as the sum of the unstained areas of the sections, multiplied by their thickness.

α_2 -Antiplasmin levels in murine plasma were measured by a chromogenic substrate assay, based on its rapid inhibition of plasmin, according to the 20 procedure described by Edy et al. in *Thromb. Res.* (1976) 8:513. Briefly, 10 μ l murine plasma (diluted 1/10 in 0.05 M NaH_2PO_4 buffer, pH 7.4, containing 0.01 percent Tween 20) was mixed at 37°C with 420 μ l 0.05 M Tris.HCl, 0.1 M NaCl buffer, pH 7.4, containing 0.01% Tween 20, and with 20 μ l of 0.125 μM human plasmin (final concentration 5 nM). After 10 seconds incubation, 50 μ l of 3 mM 25 S2403 (Chromogenix, Antwerp, Belgium) was added and the change in absorbance measured at 405 nm. Changes in absorbance of 0.18 min^{-1} for buffer and 0.09 min^{-1} for pooled murine plasma were used for the construction of a calibration curve. Data are presented in tables 2a and 2b.

EXAMPLE 5 - effect of recombinant human microplasmin on cerebral infarct size
30 in mice

Effect of recombinant microplasmin on α_2 -antiplasmin and fibrinogen levels

Effects of an intravenous bolus injection of the recombinant human microplasmin of example 2 in inbred BALB/c mice on plasma α_2 -antiplasmin and fibrinogen levels are summarized in Table 2a. The α_2 -antiplasmin and fibrinogen 5 levels decreased proportionally to the microplasmin dose and partially recovered within one hour, suggesting that α_2 -antiplasmin depletion was transient during the first few hours after the single bolus injection of microplasmin.

Effect of recombinant microplasmin on cerebral infarct size

Ligation of the MCA induced a cerebral infarct with a volume of 29 μ l in 10 inbred BALB/c mice (Table 2b). Injection of 0.07 mg recombinant human microplasmin had no significant effect on infarct size, whereas injection of 0.13 mg microplasmin or more produced a significant reduction of cerebral infarct size. This is consistent with the above transient minor reduction of α_2 -antiplasmin with the lower dose and the more persistent depletion obtained with 0.2 mg 15 microplasmin.

Table 2a

Compound	Dose (mg)	Residual		Residual	
		α_2 -antiplasmin (%)	Fibrinogen (%)	15 min.	1 h.
Saline	-	99	-	100	-
Microplasmin	0.07	58	75	93	79
	0.13	14	44	33	17
	0.20	0	18	ND	ND

ND: not determined

Table 2b

Compound	Dose (mg)	Cerebral Infarct Size (mm ³)	p*
Saline	-	29 (27-30)	-
Microplasmin	0.07	29 (27-30)	0.74
id.	0.13	26 (21-28)	0.041
Id.	0.13+0.07**	26 (20-28)	0.02

The data represent median and range of values obtained in 6 experiments

* versus saline

** injected 15' and 60' after MCA occlusion respectively

5 EXAMPLE 6 - rabbit extracorporeal loop thrombolysis model (general procedure)

A simple extracorporeal loop thrombosis model in rabbits was used for the quantitative evaluation of the thrombolytic effect of human plasmin and microplasmin, as disclosed by Hotchkiss et al. in *Thromb. Haemost.* (1987) 58: 107.

10 New Zealand white rabbits with a body weight of 2.6 – 3.2 kg were anesthetized by intramuscular injection of 1.0 of 2% xylazine and 0.5 ml of ketamine (same suppliers as in example 4). Additional Nembutal (12 mg/hour) was administered to maintain anesthesia. Thyroidal uptake of radioiodide was blocked by administration of sodium iodide (0.5 ml of a 2% solution). A femoral vein 15 catheter was introduced for blood sampling and a femoral artery catheter for blood pressure measurement (PDCR 75 from Druck Ltd, Leicester, United Kingdom).

A 300 µl thrombus was formed around a woollen thread introduced longitudinally in each of two adapted insulin syringes from a mixture of ¹²⁵I-labeled fibrinogen (approximately 400,000 cpm), platelet poor rabbit plasma, and 0.07 ml

thrombin solution (100 NIH U/ml). In all instances, the clot formed quickly and was allowed to age for 30 minutes at 37°C. The two syringes were then inserted in an extracorporeal loop of silicon tubing between a femoral artery and a marginal ear vein. The blood flow was regulated via a peristaltic pump (P1 available from 5 Pharmacia LKB, Piscataway, New Jersey). Thrombotic extension of the clot was prevented by infusion of heparin (300 U/kg bolus followed by 200 U/kg over 2 hours) and the platelet aggregation inhibitor Ridogrel (7.5 mg/kg) bolus, 30 minutes before starting infusion of wild-type plasmin (available from Janssen Research Foundation, Beerse, Belgium) or the recombinant microplasmin obtained in 10 example 2. The extent of thrombolysis was measured as the difference between the radioactivity introduced in the clot and that recovered in the syringes at the end of the experiment.

Local infusion was carried out by using a constant rate infusion pump (Perfuser VI, available from B. Braun, Penang, Malaysia), through a three-ways 15 valve, in a volume of 6 ml over 2 hours proximal to the first inserted syringe in the extracorporeal loop. The extent of thrombolysis was calculated 2.5 hours after starting infusion, as the difference between the radioactivity originally incorporated in the clot and the radioactivity in the syringe, and expressed as a percentage of the initial radioactivity.

20 2-ml blood samples were drawn into trisodium citrate (final concentration 0.011 M) before starting infusion and at hourly intervals for 2 hours. These samples were used for measurements of fibrinogen, α_2 -antiplasmin, and activated partial thromboplastin time. Bleeding times were performed by applying a Symplate II device (available from Organon Technica, Durham, North Carolina) to 25 a shaved inner thigh surface.

EXAMPLE 7 - effect of recombinant microplasmin on extracorporeal loop clot lysis

Results of the determinations made in accordance with the general procedure of example 6 are presented in the following Table 3. Clot lysis with the recombinant microplasmin of example 2 produced minor α_2 -antiplasmin depletion

and fibrinogen breakdown and was associated with minor bleeding time prolongation. Infusion of wild-type plasmin resulted in a reduction of 80% of α_2 -antiplasmin and fibrinogen levels, with minor effect on the bleeding time. These findings indicate that the extent of clot lysis by recombinant microplasmin and wild-type plasmin is mainly determined by the dose of the drug and its delivery in the vicinity of the thrombus. Thrombolysis with recombinant microplasmin or wild-type plasmin thus was not associated with extensive systemic activation of the fibrinolytic system as evidenced by the moderate changes in fibrinogen, α_2 -antiplasmin and bleeding time.

10

Table 3

Substance	Blood flow (ml/min)	Dose (mg/kg)	Clot Lysis (percent)	Residual Fibrinogen (percent)*	Residual α_2 -antiplasmin (percent)*	Bleeding time (sec)*
Solvent	-	-	26	110	110	90
Microplasmin	0.5	0.6	43	93	97	95
		1.3	46	93	96	145
		2.5	51	79	71	120
		3.8	64	77	71	150
		5.0	60	67	67	90
	0.1	2.5	80	87	87	98
Plasmin	0.5	2.5	44	82	80	170
		5.0	53	80	86	170
	0.1	2.5	56	93		120

*: at the end of the infusion.

EXAMPLE 8 - dog circumflex coronary artery copper coil-induced thrombosis (general procedure)

A copper coil was introduced in the coronary artery for the quantitative evaluation of the thrombolytic effect of human wild-type plasmin and recombinant microplasmin as described by Bergmann et al. in *Science* (1983) 220:1181-1183. Dogs were anesthetized by intravenous injection of 30 mg/kg Nembutal of after premedication by intramuscular injection with atropine 12.5 μ g/kg and ketamine 10

mg/kg (same suppliers as in example 4). Anesthesia was maintained by Nembutal infusion of 8 to 10 mg/kg/hour. After pre-medication, an intravenous line is introduced in the front leg vein and fix. This intravenous line is taken for anaesthetic drug administration and infusion. A second venous access line is 5 introduced in a saphenous vein for heparin administration. During further preparation the line is kept open with a saline infusion at approx. 20 ml/hour. The femoral artery is exposed via an incision close to the groin and a catheter is introduced for blood sampling. The right and left carotid arteries are exposed for coronary catheterisation. ECG electrodes and a rectal temperature probe are 10 placed for continuous monitoring of the ECG and the body temperature. The copper coil should be rinsed with 50% acetic acid to remove oxidation to have an optimal thrombogenic copper surface. A Lehman catheter (5 Fg, USCI Bard) is connected to the angiographic valve system introduced into the left carotid artery and advanced into the left coronary artery. An angiogram is performed to identify 15 the position of the first major side branch of the circumflex artery. The angiographic valve system is disconnected and thin guide wire is introduced through the Lehman catheter and positioned distally to the first dominant side branch. The angiographic valve system should have the following connections: contrast medium, pressurized saline, pressure transducer (to monitor non occlusive catheter placement) and 20 infusion line for intracoronary recombinant microplasmin administration. The Lehman catheter is retracted and removed, while the guidewire is kept in position. Over this guide wire the copper coil, fixed to a second guidewire is introduced and advanced into the circumflex artery and placed distally to the first dominant side branch. A selection of copper coils is available and the best estimated size used 25 e.g. a 3-mm copper coil (six turns of a 0.5 mm copper wire). The coil must be distal from the predominant side branch if not fibrillation upon occlusion might occur. The central guide wire is removed. Via the right carotid artery the Lehman catheter is reintroduced to perform the angiograms throughout the remaining part of the study. The formation of a thrombotic occlusion is monitored by 30 electrocardiographic measurements or by angiography via the right carotid artery. The occlusion was confirmed by angiography. The occlusion is aged for 60

minutes. An angiogram is then performed to confirm total occlusion of the artery and the Lehman catheter is left in place, proximal to the thrombus without occlusion of the circumflex coronary artery for administration of recombinant microplasmin and to perform the angiograms. Then a heparin 200 units/kg 5 intravenous bolus followed by a 40 units/kg/hour infusion at 1 ml/min intravenous infusion throughout the experiment is given via the saphenous vein catheter. Five minutes after heparin bolus administration, a first bolus over 5 minutes of 40 % of the recombinant microplasmin dose is administered intracoronary via the Lehman catheter. If occlusion persists after 15 minutes, as evidenced by angiography, the 10 remaining 60 % of the recombinant microplasmin dose will be infused intracoronary over 1 hour. Reperfusion and reocclusion are evaluated angiographically at 15 minutes intervals or whenever electrocardiographic signs suggestive for reperfusion or re-occlusion occurred. At the end of the experiment, the animals are killed by administering an overdose of pentobarbital (10 ml of 60 mg/ml 15 intravenously). Six blood samples of 4.5 ml blood are taken 0.5 ml 3.8% trisodium citrate (final concentration 0.011 M) and kept on ice. The timing of the blood samples was baseline (after 45 minutes of stable occlusion, 2 minutes after heparin bolus but before recombinant microplasmin administration, 5 minutes after 20 recombinant microplasmin bolus administration, at end of infusion and 120 minutes after the end of the infusion). These samples are centrifugated at 2000 rpm for 10 minutes at 4°C . The plasma is collected and frozen at -20°C for determination of fibrinogen and α_2 -antiplasmin according to Edy et al.(cited *supra*).

EXAMPLE 9 - effect of recombinant microplasminogen on copper-coil induced myocardial infarction

25 Results of the determinations made in accordance with the general procedure of example 8 are presented in table 4 and figure 2. An erythrocyte-rich thrombus was formed within 15 minutes after introduction of the copper coil as evidenced electrocardiographic signs induced by the transmural ischemia and confirmed angiography via the right carotid artery. Two doses of the recombinant 30 microplasminogen of example 2 were investigated in four animals per group. The

first group received a bolus of 2 mg/kg over 5 minutes and, if occlusion persisted 15 minutes later as evidenced by angiography, a residual dose of 3 mg/kg was initiated over one hour. The second group received a bolus of 1 mg/kg and, if occlusion persisted 15 minutes later as evidenced by angiography, infusion of the 5 residual 1.5 mg/kg was initiated over one hour.

In the first group, three dogs treated with the 2 mg/kg had a complete and persistent resolution of the thrombus within 15 minutes after administration. In the fourth dog, a complete and persistent partial recanalisation occurred after administration of the full dose (figure). In the second group, one dog had a 10 complete and persistent reperfusion 15 minutes after bolus injection. In the remaining three dogs, the bolus followed by a 1 hour infusion of 1.5 m/kg induced a complete and persistent recanalisation (figure).

As shown in table 4, administration of recombinant microplasminogen induced only a partial decrease of fibrinogen and α_2 -antiplasmin.

15

Table 4

Total Dose (mg/kg)	α_2 -antiplasmin (%)				Fibrinogen (g/l)			
	Baseline	After heparin	15 min after bolus	End of experiments	Baseline	After heparin	15 min after bolus	End of experiments
2	105	97	45	56	1.43	1.40	0.73	0.74
2.5	106	105	83	58	1.78	1.88	1.37	1.03

CLAIMS

1. A yeast expression vector comprising a mammalian nucleotide sequence operably linked to a promoter, wherein the said mammalian nucleotide sequence codes for the catalytic domain of plasminogen and further optionally codes for one or more kringle domains of plasminogen, or mutants or hybrids thereof.
5
2. The yeast expression vector of claim 1 wherein the mammalian nucleotide sequence codes for plasminogen.
10
3. The yeast expression vector of claim 1 wherein the mammalian nucleotide sequence codes for microplasminogen.
4. The yeast expression vector of claim 1 wherein the mammalian nucleotide sequence codes for miniplasminogen.
15
5. A yeast expression vector according to any of claims 1 to 4, wherein the nucleotide sequence is a human nucleotide sequence.
- 20 6. The yeast expression vector of claim 1, wherein the nucleotide sequence is SEQ ID No.1.
7. The yeast expression vector of claim 1, where the nucleotide sequence is SEQ. ID. No. 3.
25
8. The yeast expression vector of claim 1, wherein the nucleotide sequence is SEQ. ID. No.5.
9. A yeast expression vector according to any of claims 1 to 8, wherein the promoter is an inducible promoter.
30

10. The yeast expression vector of any of claims 1 to 9, which is able to stably integrate in the yeast genome, e.g. by homologous recombination.
11. The yeast expression vector of any of claims 1 to 9, wherein the nucleotide sequence is fused to a secretion signal.
12. The yeast expression vector of any of claims 1 to 11, for expression in a yeast selected from the group consisting of *Hansenula*, *Pichia*, *Candida* and *Torulopsis genera*.

10

13. A yeast cell comprising a mammalian nucleotide sequence coding for the catalytic domain of plasminogen and further optionally codes for one or more kringle domains of plasminogen, or mutants or hybrids thereof.

15

14. A yeast cell according to claim 13, wherein the said mammalian nucleotide sequence codes for plasminogen.
15. A yeast cell according to claim 13, wherein the said mammalian nucleotide sequence codes for miniplasminogen.

20

16. A yeast cell according to claim 13, wherein the said mammalian nucleotide sequence codes for microplasminogen.

17. A yeast cell comprising the nucleotide sequence SEQ.ID.No. 1.

25

18. A yeast cell comprising the nucleotide sequence SEQ.ID.No. 3.
19. A yeast cell comprising the nucleotide sequence SEQ.ID.No. 5.

30

20. A yeast cell transfected with a vector according to any of claims 1 to 12.

21. A yeast cell according to any of claims 13 to 20, belonging to the group of the methylotrophic yeasts.

22. A yeast cell according to any of claims 13 to 21, wherein the said yeast is
5 selected from the group consisting of the, *Hansenula*, , *Pichia*, *Candida* and
Torulopsis genera.

23. A yeast cell according to any of claims 13 to 20, belonging to the *Pichia pastoris* species.

10

24. A yeast cell according to any of claims 20 to 23, wherein the said vector is integrated into the genome.

15

25. A yeast cell according to any of claims 13 to 24, belonging to the deposited cell line with accession number MUCL43676.

26. A yeast cell according to any of claims 13 to 25, being able to express human microplasminogen at a level of at least about 100 mg/litre.

20

27. A yeast cell according to any of claims 13 to 25, being able to express human miniplasminogen at a level of at least about 3 mg/litre.

25

28. A method of expressing a mammalian protein comprising the catalytic domain of plasminogen and further optionally comprising one or more kringle domains of plasminogen, or mutants or hybrids thereof, in a yeast cell according to any of claims 13 to 25, using recombinant technology procedures.

29. A method according to claim 28, wherein the mammalian protein has the amino-acid sequence SEQ.ID. No. 2.

30

30. A method according to claim 28, wherein the mammalian protein has the amino-acid sequence SEQ.ID. No. 4.

5 31. A method according to claim 28, wherein the mammalian protein has the amino-acid sequence SEQ.ID. No. 6.

32. A method according to any of claims 28 to 31, further comprising the step of activating the expressed mammalian protein by means of a plasminogen activator.

10

33. A method according to claim 32, wherein the said plasminogen activator is staphylokinase or a variant thereof.

15

34. A method according to claim 32 or claim 33, further comprising the step of stabilizing the expressed and activated mammalian protein by means of a stabilizing agent.

20

35. A method according to claim 34, wherein the said stabilizing agent comprises an amino-acid selected from the group consisting of lysine, 6-amino hexanoic acid and tranexamic acid.

36. A method according to claim 34, wherein the said stabilizing agent comprises a stabilizing medium.

25

37. A method according to claim 34, wherein the said stabilizing medium is an acid solution or an acid buffer.

38. A method according to claim 36 or claim 37, wherein the said stabilizing medium is a citrate buffer with a pH of about 3.1.

30

39. A method according to any of claims 34 to 38, further comprising the step of drying the expressed, activated and stabilized mammalian protein by means of lyophilization.

5 40. A recombinant mammalian protein obtained by a method according to any of claim 28 to 39 or expressed in a yeast cell according to any of claims 1 to 27.

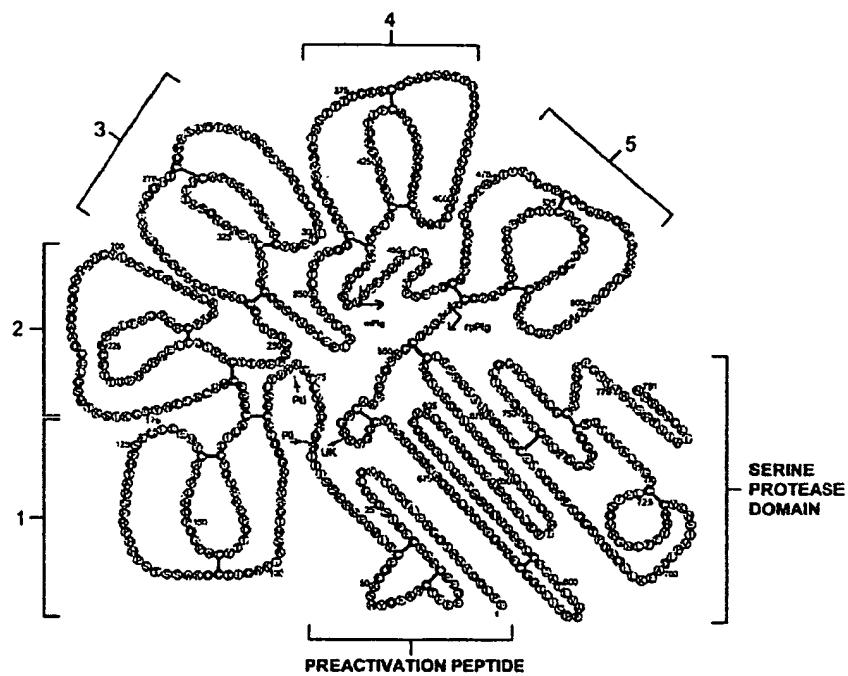
10 41. A method of treatment of a thromboembolic disease in a mammal, comprising administration to the said mammal an effective amount of the recombinant mammalian protein of claim 40.

15

20

25

1/5



1 - 5 : KRINGLE DOMAINS

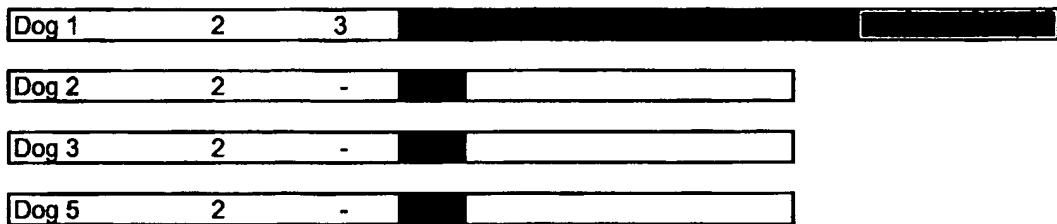
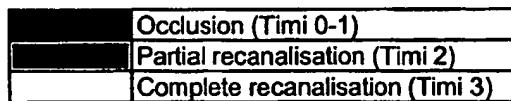
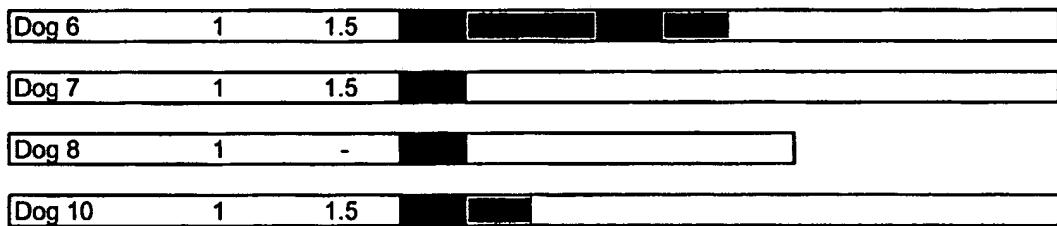
Figure 1

2 / 5

Figure 2

5

	bolus (mg/kg)	infusion (mg/kg)	0	15	30	45	60	75	90	105	120	135
--	------------------	---------------------	---	----	----	----	----	----	----	-----	-----	-----

Dose Group I**Dose Group II**

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FIGURE 3

10	20	30	40	
GCC CCT TCA TTT GAT TGT GGG AAG CCT CAA GTG GAG CCG AAG AAA TGT				
A P S F D C G K P Q V E P K K C>				
50	60	70	80	90
CCT GGA AGG GTT GTA GGG GGG TGT GTG GCC CAC CCA CAT TCC TGG CCC				
P G R V V G G C V A H P H S W P>				
100	110	120	130	140
TGG CAA GTC AGT CTT AGA ACA AGG TTT GGA ATG CAC TTC TGT GGA GGC				
W Q V S L R T R F G M H F C G G>				
150	160	170	180	190
ACC TTG ATA TCC CCA GAG TGG GTG TTG ACT GCA GCC CAC TGC TTG GAG				
T L I S P E W V L T A A A H C L E>				
200	210	220	230	240
AAG TCC CCA AGG CCT TCA TCC TAC AAG GTC ATC CTA GGT GCA CAC CAA				
K S P R P S S Y K V I L G A H Q>				
250	260	270	280	
GAA GTG AAT CTC GAA CCG CAT GTT CAG GAA ATA GAA GTG TCT AGG CTG				
E V N L E P H V Q E I E V S R L>				
290	300	310	320	330
TTC TTG GAG CCC ACA CGA AAA GAT ATT GCC TTG CTA AAG CTA AGC AGT				
F L E P T R K D I A L L 'K L S S>				
340	350	360	370	380
CCT GCC GTC ATC ACT GAC AAA GTA ATC CCA GCT TGT CTG CCA TCC CCA				
P A V I T D K V I P A C L P S P>				
390	400	410	420	430
AAT TAT GTG GTC GCC GAC CGG ACC GAA TGT TTC ATC ACT GGC TGG GGA				
N Y V V A D R T E C F I T G W G>				
440	450	460	470	480
GAA ACC CAA GGT ACT TTT GGA GCT GGC CTT CTC AAG GAA GCC CAG CTC				
E T Q G T F G A G L L K E A Q L>				
490	500	510	520	
CCT GTG ATT GAG AAT AAA GTG TGC AAT CGC TAT GAG TTT CTG AAT GGA				
P V I E N K V C N R Y E F L N G>				
530	540	550	560	570
AGA GTC CAA TCC ACC GAG CTC TGT GCT GGG CAT TTG GCC GGA GGC ACT				
R V Q S T E L C A G H L A G G T>				
580	590	600	610	620
GAC AGT TGC CAG GGT GAC AGT GGA GGG CCT CTG GTT TGC TTC GAG AAG				
D S C Q G D S G G P L V C F E K>				
630	640	650	660	670
GAC AAA TAC ATT TTA CAA GGA GTC ACT AGT TGG GGT CTT GGC TGT GCA				
D K Y I L Q G V T S W G L G C A>				
680	690	700	710	720
CGC CCC AAT AAG CCT GGT GTC TAT GTT CGT GTC TCC AGG TTT GTT ACT				
R P N K P G V Y V R V S R F V T>				
730	740	750		
TGG ATT GAG GGA GTG ATG AGA AAT AAT TAA				
W I E G V M R N N>				

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FIGURE 4

10 20 30 40 50 60
 GCA CCT CCG CCT GTT GTC CTG CTT CCA GAT GTA GAG ACT CCT TCC GAA GAA GAC TGT ATG TTT
 GGG AAT
 A P P P V V L L P D V E T P S E E D C M F G
 N

70 80 90 100 110 120 130
 GGG AAA GGA TAC CGA GGC AAG AGG GCG ACC ACT GTT ACT GGG ACG CCA TGC CAG GAC TGG GCT
 GCC CAG
 G K G Y R G K R A T T V T G T P C Q D A W
 A Q

140 150 160 170 180 190 200
 GAG CCC CAT AGA CAC AGC ATT TTC ACT CCA GAG ACA AAT CCA CGG GCG GGT CTG GAA AAA AAT
 TAC TGC
 E P H R H S I F T P E T N P R A G L E K N
 Y C

210 220 230 240 250 260 270
 CGT AAC CCT GAT GGT GAT GTA GGT CCC TGG TGC TAC ACG ACA AAT CCA AGA AAA CTT TAC
 GAC TAC
 R N P D G D V G G P W C Y T T N P R K L Y D
 Y

280 290 300 310 320 330 340
 TGT GAT GTC CCT CAG TGT GCG GCC CCT TCA TTT GAT TGT GGG AAG CCT CAA GTG GAG CCG AAG
 AAA TGT
 C D V P Q C A A P S F D C G K P Q V E P K K
 C

350 360 370 380 390 400 410
 CCT GGA AGG GTT GTA GGG GGG TGT GTG GCC CAC CCA CAT TCC TGG CCC TGG CAA GTC AGT CTT
 AGA ACA
 P G R V V G G C V A H P H S W P W Q V S L R
 T

420 430 440 450 460 470
 480 490 500 510 520 530 540
 AGG TTT GGA ATG CAC TTC TGT GGA GGC ACC TTG ATA TCC CCA GAG TGG GTG TTG ACT GCA GCC
 CAC TGC
 R F G M H F C G G T L I S P E W V L T A A H
 C

550 560 570 580 590 600 610
 TTG GAG AAG TCC CCA AGG CCT TCA TCC TAC AAG GTC ATC CTA GGT GCA CAC CAA GAA GTG AAT
 CTC GAA
 L E K S P R P S S Y K V I L G A H Q E V N L
 E

620 630 640 650 660 670 680
 CCG CAT GTT CAG GAA ATA GAA GTG TCT AGG CTG TTC TTG GAG CCC ACA CGA AAA GAT ATT GCC
 TTG CTA
 P H V Q E I E V S R L F L E P T R K D I A L
 L

690 700 710 720 730 740 750
 AAG CTA AGC AGT CCT GCC GTC ATC ACT GAC AAA GTA ATC CCA GCT TGT CTG CCA TCC CCA AAT
 TAT GTG
 K L S S P A V I T D K V I P A C L P S P N Y
 V

760 770 780 790 800 810 820

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Figure 4 (continuation)

CTC AAG GAA GCC CAG CTC CCT GTG ATT GAG AAT AAA GTG TGC AAT CGC TAT GAG TTT CTG AAT
 GGA AGA
 L K E A Q L P V I E N K V C N R Y E F L N
 G R

830 840 850 860 870 880 890
 GTC CAA TCC ACC GAG CTC TGT GCT GGG CAT TTG GCC GGA GGC ACT GAC AGT TGC CAG GGT GAC
 AGT GGA
 V Q S T E L C A G H L A G G T D S C Q G D
 S G

900 910 920 930 940 950 960
 GGG CCT CTG GTT TGC TTC GAG AAG GAC AAA TAC ATT TTA CAA GGA GTC ACT AGT TGG GGT CTT
 GGC TGT
 G P L V C F E K D K Y I L Q G V T S W G L G
 C

970 980 990 1000 1010 1020 1030
 GCA CGC CCC AAT AAG CCT GGT GTC TAT GTT CGT GTC TCC AGG TTT GTT ACT TGG ATT GAG GGA
 GTG ATG
 A R P N K P G V Y V R V S R F V T W I E G V
 M

1040
 AGA AAT AAT TAA
 R N N>

SEQUENCE LISTING

<110> Thromb-X N.V.
Collen, Désiré
Laroche, Yves
Naqai, Nubuo

<120> A yeast expression system vector and a method of making a recombinant protein by expression in a yeast cell.

<130> T-1923 PCT

<150> gb2000200031196
<151> 2000-12-21

<150> gb2001200116690
<151> 2001-07-09

<150> gb2001200116702
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Met Glu His Lys Glu Val Val Leu Leu Leu Leu Leu Phe Leu Lys Ser
1 5 10 15

48

```

ggc caa gga gag cct ctg gat gac tat gtg aat acc cag ggg gct tca
Gly Gln Gly Glu Pro Leu Asp Asp Tyr Val Asn Thr Gln Gly Ala Ser
20          25          30

```

96

ctg ttc agt gtc act aag aag cag ctg gga gca gga agt ata gaa gaa
 Leu Phe Ser Val Thr Lys Lys Gln Leu Gly Ala Gly Ser Ile Glu Glu
 35 40 45

144

tgt	gca	gca	aaa	tgt	gag	gag	gac	gaa	gaa	ttc	acc	tgc	agg	gca	ttc
Cys	Ala	Ala	Lys	Cys	Glu	Glu	Asp	Glu	Glu	Phe	Thr	Cys	Arg	Ala	Phe
50	55									60					

192

caa	tat	cac	agt	aaa	gag	caa	caa	tgt	gtg	ata	atg	gct	gaa	aac	agg
Gln	Tyr	His	Ser	Lys	Glu	Gln	Gln	Cys	Val	Ile	Met	Ala	Glu	Asn	Arg
65		70						75						80	

240

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aag tcc tcc ata atc att agg atg aga gat gta gtt tta ttt gaa aag
Lys Ser Ser Ile Ile Arg Met Arg Asp Val Val Val Leu Phe Glu Lys
          85           90           95

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288

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ggg acg atg tcc aaa aca aaa aat ggc atc acc tgtcaa aaa tgg agt Gly Thr Met Ser Lys Thr Lys Asn Gly Ile Thr Cys Gln Lys Trp Ser 115 120 125	384
tcc act tct ccc cac aga cct aga ttc tca cct gct aca cac ccc tca Ser Thr Ser Pro His Arg Pro Arg Phe Ser Pro Ala Thr His Pro Ser 130 135 140	432
gag gga ctg gag gag aac tac tgc agg aat cca gac aac gat ccg cag Glu Gly Leu Glu Glu Asn Tyr Cys Arg Asn Pro Asp Asn Asp Pro Gln 145 150 155 160	480
ggg ccc tgg tgc tat act act gat cca gaa aag aga tat gac tac tgc Gly Pro Trp Cys Tyr Thr Asp Pro Glu Lys Arg Tyr Asp Tyr Cys 165 170 175	528
gac att ctt gag tgt gaa gag gaa tgt atg cat tgc agt gga gaa aac Asp Ile Leu Glu Cys Glu Glu Cys Met His Cys Ser Gly Glu Asn 180 185 190	576
tat gac ggc aaa att tcc aag acc atg tct gga ctg gaa tgc cag gcc Tyr Asp Gly Ile Ser Lys Thr Met Ser Gly Leu Glu Cys Gln Ala 195 200 205	624
tgg gac tct cag agc cca cac gct cat gga tac att cct tcc aaa ttt Trp Asp Ser Gln Ser Pro His Ala His Gly Tyr Ile Pro Ser Lys Phe 210 215 220	672
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tgc gac atc ccc cgc tgc aca aca cct cca cca tct tct ggt ccc acc Cys Asp Ile Pro Arg Cys Thr Thr Pro Pro Pro Ser Ser Gly Pro Thr 260 265 270	816
tac cag tgt ctg aag gga aca ggt gaa aac tat cgc ggg aat gtg gct Tyr Gln Cys Leu Lys Gly Thr Gly Glu Asn Tyr Arg Gly Asn Val Ala 275 280 285	864
gtt acc gtt tcc ggg cac acc tgt cag cac tgg agt gca cag acc cct Val Thr Val Ser Gly His Thr Cys Gln His Trp Ser Ala Gln Thr Pro 290 295 300	912
cac aca cat aac agg aca cca gaa aac ttc ccc tgc aaa aat ttg gat His Thr His Asn Arg Thr Pro Glu Asn Phe Pro Cys Lys Asn Leu Asp 305 310 315 320	960
gaa aac tac tgc cgc aat cct gac gga aaa agg gcc cca tgg tgc cat Glu Asn Tyr Cys Arg Asn Pro Asp Gly Lys Arg Ala Pro Trp Cys His 325 330 335	1008
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Asp Ser Ser Pro Val Ser Thr Glu Gln Leu Ala Pro Thr Ala Pro Pro			
355	360	365	
gag cta acc cct gtg gtc cag gac tgc tac cat ggt gat gga cag agc			1152
Glu Leu Thr Pro Val Val Gln Asp Cys Tyr His Gly Asp Gly Gln Ser			
370	375	380	
tac cga ggc aca tcc tcc acc acc aca gga aag aag tgt cag tct			1200
Tyr Arg Gly Thr Ser Ser Thr Thr Thr Gly Lys Lys Cys Gln Ser			
385	390	395	400
tgg tca tct atg aca cca cac cgg cac cag aag acc cca gaa aac tac			1248
Trp Ser Ser Met Thr Pro His Arg His Gln Lys Thr Pro Glu Asn Tyr			
405	410	415	
cca aat gct ggc ctg aca atg aac tac tgc agg aat cca gat gcc gat			1296
Pro Asn Ala Gly Leu Thr Met Asn Tyr Cys Arg Asn Pro Asp Ala Asp			
420	425	430	
aaa ggc ccc tgg tgt ttt acc aca gac ccc agc gtc agg tgg gag tac			1344
Lys Gly Pro Trp Cys Phe Thr Thr Asp Pro Ser Val Arg Trp Glu Tyr			
435	440	445	
tgc aac ctg aaa aaa tgc tca gga aca gaa gcg agt gtt gta gca cct			1392
Cys Asn Leu Lys Lys Cys Ser Gly Thr Glu Ala Ser Val Val Ala Pro			
450	455	460	
ccg cct gtt gtc ctg ctt cca gat gta gag act cct tcc gaa gaa gac			1440
Pro Pro Val Val Leu Leu Pro Asp Val Glu Thr Pro Ser Glu Glu Asp			
465	470	475	480
tgt atg ttt ggg aat ggg aaa gga tac cga ggc aag agg gcg acc act			1488
Cys Met Phe Gly Asn Gly Lys Gly Tyr Arg Gly Lys Arg Ala Thr Thr			
485	490	495	
gtt act ggg acg cca tgc cag gac tgg gct gcc cag gag ccc cat aga			1536
Val Thr Gly Thr Pro Cys Gln Asp Trp Ala Ala Gln Glu Pro His Arg			
500	505	510	
cac agc att ttc act cca gag aca aat cca cgg gcg ggt ctg gaa aaa			1584
His Ser Ile Phe Thr Pro Glu Thr Asn Pro Arg Ala Gly Leu Glu Lys			
515	520	525	
aat tac tgc cgt aac cct gat ggt gat gta ggt ggt ccc tgg tgc tac			1632
Asn Tyr Cys Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Cys Tyr			
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acg aca aat cca aga aaa ctt tac gac tac tgt gat gtc cct cag tgt			1680
Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr Cys Asp Val Pro Gln Cys			
545	550	555	560
gcg gcc cct tca ttt gat tgt ggg aag cct caa gtg gag ccg aag aaa			1728
Ala Ala Pro Ser Phe Asp Cys Gly Lys Pro Gln Val Glu Pro Lys Lys			
565	570	575	
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Cys Pro Gly Arg Val Val Gly Gly Cys Val Ala His Pro His Ser Trp			

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595	600	605	
ggc acc ttg ata tcc cca gag tgg gtg ttg act gct gcc cac tgc ttg Gly Thr Leu Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His Cys Leu	610	615	1872
610	615	620	
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675	680	685	
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690	695	700	
gga gaa acc caa ggt act ttt gga gct ggc ctt ctc aag gaa gcc cag Gly Glu Thr Gln Gly Thr Phe Gly Ala Gly Leu Leu Lys Glu Ala Gln	705	710	2160
705	710	715	
715	720		
ctc cct gtg att gag aat aaa gtg tgc aat cgc tat gag ttt ctg aat Leu Pro Val Ile Glu Asn Lys Val Cys Asn Arg Tyr Glu Phe Leu Asn	725	730	2208
725	730	735	
gga aga gtc caa tcc acc gaa ctc tgt gct ggg cat ttg gcc gga ggc Gly Arg Val Gln Ser Thr Glu Leu Cys Ala Gly His Leu Ala Gly	740	745	2256
740	745	750	
750			
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755	760	765	
aag gac aaa tac att tta caa gga gtc act tct tgg ggt ctt ggc tgt Lys Asp Lys Tyr Ile Leu Gln Gly Val Thr Ser Trp Gly Leu Gly Cys	770	775	2352
770	775	780	
780			
gca cgc ccc aat aag cct ggt gtc tat gtt cgt gtt tca agg ttt gtt Ala Arg Pro Asn Lys Pro Gly Val Tyr Val Arg Val Ser Arg Phe Val	785	790	2400
785	790	795	
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20 25 30

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35 40 45

Cys Ala Ala Lys Cys Glu Glu Asp Glu Glu Phe Thr Cys Arg Ala Phe
50 55 60

Gln Tyr His Ser Lys Glu Gln Gln Cys Val Ile Met Ala Glu Asn Arg
65 70 75 80

Lys Ser Ser Ile Ile Ile Arg Met Arg Asp Val Val Leu Phe Glu Lys
85 90 95

Lys Val Tyr Leu Ser Glu Cys Lys Thr Gly Asn Gly Lys Asn Tyr Arg
100 105 110

Gly Thr Met Ser Lys Thr Lys Asn Gly Ile Thr Cys Gln Lys Trp Ser
115 120 125

Ser Thr Ser Pro His Arg Pro Arg Phe Ser Pro Ala Thr His Pro Ser
130 135 140

Glu Gly Leu Glu Glu Asn Tyr Cys Arg Asn Pro Asp Asn Asp Pro Gln
145 150 155 160

Gly Pro Trp Cys Tyr Thr Thr Asp Pro Glu Lys Arg Tyr Asp Tyr Cys
165 170 175

Asp Ile Leu Glu Cys Glu Glu Cys Met His Cys Ser Gly Glu Asn
180 185 190

Tyr Asp Gly Lys Ile Ser Lys Thr Met Ser Gly Leu Glu Cys Gln Ala
195 200 205

Trp Asp Ser Gln Ser Pro His Ala His Gly Tyr Ile Pro Ser Lys Phe
210 215 220

Pro Asn Lys Asn Leu Lys Lys Asn Tyr Cys Arg Asn Pro Asp Arg Glu
225 230 235 240

Leu Arg Pro Trp Cys Phe Thr Thr Asp Pro Asn Lys Arg Trp Glu Leu
245 250 255

Cys Asp Ile Pro Arg Cys Thr Thr Pro Pro Pro Ser Ser Gly Pro Thr
260 265 270

Tyr Gln Cys Leu Lys Gly Thr Gly Glu Asn Tyr Arg Gly Asn Val Ala
275 280 285

Val Thr Val Ser Gly His Thr Cys Gln His Trp Ser Ala Gln Thr Pro
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His Thr His Asn Arg Thr Pro Glu Asn Phe Pro Cys Lys Asn Leu Asp
305 310 315 320

Glu Asn Tyr Cys Arg Asn Pro Asp Gly Lys Arg Ala Pro Trp Cys His
325 330 335

Thr Thr Asn Ser Gln Val Arg Trp Glu Tyr Cys Lys Ile Pro Ser Cys
340 345 350

Asp Ser Ser Pro Val Ser Thr Glu Gln Leu Ala Pro Thr Ala Pro Pro
355 360 365

Glu Leu Thr Pro Val Val Gln Asp Cys Tyr His Gly Asp Gly Gln Ser
370 375 380

Tyr Arg Gly Thr Ser Ser Thr Thr Thr Gly Lys Lys Cys Gln Ser
385 390 395 400

Trp Ser Ser Met Thr Pro His Arg His Gln Lys Thr Pro Glu Asn Tyr
405 410 415

Pro Asn Ala Gly Leu Thr Met Asn Tyr Cys Arg Asn Pro Asp Ala Asp
420 425 430

Lys Gly Pro Trp Cys Phe Thr Thr Asp Pro Ser Val Arg Trp Glu Tyr
435 440 445

Cys Asn Leu Lys Lys Cys Ser Gly Thr Glu Ala Ser Val Val Ala Pro
450 455 460

Pro Pro Val Val Leu Leu Pro Asp Val Glu Thr Pro Ser Glu Glu Asp

465

470

475

480

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Val Thr Gly Thr Pro Cys Gln Asp Trp Ala Ala Gln Glu Pro His Arg
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Pro Trp Gln Val Ser Leu Arg Thr Arg Phe Gly Met His Phe Cys Gly
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Gly Thr Leu Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His Cys Leu
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Glu Lys Ser Pro Arg Pro Ser Ser Tyr Lys Val Ile Leu Gly Ala His
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Gln Glu Val Asn Leu Glu Pro His Val Gln Glu Ile Glu Val Ser Arg
645 650 655

Leu Phe Leu Glu Pro Thr Arg Lys Asp Ile Ala Leu Leu Lys Leu Ser
660 665 670

Ser Pro Ala Val Ile Thr Asp Lys Val Ile Pro Ala Cys Leu Pro Ser
675 680 685

Pro Asn Tyr Val Val Ala Asp Arg Thr Glu Cys Phe Ile Thr Gly Trp
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 Trp Gln Val Ser Leu Arg Thr Arg Phe Gly Met His Phe Cys Gly Gly
 35 40 45

acc ttg ata tcc cca gag tgg gtg act gct gcc cac tgc ttg gag 192
 Thr Leu Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His Cys Leu Glu
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 Lys Ser Pro Arg Pro Ser Ser Tyr Lys Val Ile Leu Gly Ala His Gln
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gaa gtg aat ctc gaa ccg cat gtt cag gaa ata gaa gtg tct agg ctg 288
 Glu Val Asn Leu Glu Pro His Val Gln Glu Ile Glu Val Ser Arg Leu
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aga gtc caa tcc acc gaa ctc tgt gct ggg cat ttg gcc gga ggc act Arg Val Gln Ser Thr Glu Leu Cys Ala Gly His Leu Ala Gly Thr 180 185 190	576
gac agt tgc cag ggt gac agt gga ggt cct ctg gtt tgc ttc gag aag Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Phe Glu Lys 195 200 205	624
gac aaa tac att tta caa gga gtc act tct tgg ggt ctt ggc tgt gca Asp Lys Tyr Ile Leu Gln Gly Val Thr Ser Trp Gly Leu Gly Cys Ala 210 215 220	672
cgc ccc aat aag cct ggt gtc tat gtt cgt gtt tca agg ttt gtt act Arg Pro Asn Lys Pro Gly Val Tyr Val Arg Val Ser Arg Phe Val Thr 225 230 235 240	720
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Glu Val Asn Leu Glu Pro His Val Gln Glu Ile Glu Val Ser Arg Leu
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Asn Tyr Val Val Ala Asp Arg Thr Glu Cys Phe Ile Thr Gly Trp Gly
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Arg Val Gln Ser Thr Glu Leu Cys Ala Gly His Leu Ala Gly Gly Thr
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Cys Tyr Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr Cys Asp Val Pro		
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Leu Ser Ser Pro Ala Val Ile Thr Asp Lys Val Ile Pro Ala Cys Leu		
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Ala Gln Leu Pro Val Ile Glu Asn Lys Val Cys Asn Arg Tyr Glu Phe
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Gly Gly Thr Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys
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Phe Glu Lys Asp Lys Tyr Ile Leu Gln Gly Val Thr Ser Trp Gly Leu
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INTERNATIONAL SEARCH REPORT

Inte...al Application No
PCT/BE 01/00217A. CLASSIFICATION OF SUBJECT MATTER
IPC 7 C12N15/81 C12N9/68 C12N5/10

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 7 C12N

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

SEQUENCE SEARCH, MEDLINE, BIOSIS, EPO-Internal, WPI Data, PAJ

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 93 07893 A (ORION THERAPEUTIC SYST INC) 29 April 1993 (1993-04-29) cited in the application page 16, line 17 -page 21, line 5 page 4, line 2-11 ---	40,41
X	WO 00 18436 A (LEUVEN RES & DEV VZW ;COLLEN DESIRE JOSE (GB); NOBUO NAGIA (JP)) 6 April 2000 (2000-04-06) cited in the application page 8, paragraph 2.3 example 5 ---	40,41 -/-

 Further documents are listed in the continuation of box C. Patent family members are listed in annex.

* Special categories of cited documents:

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- *O* document referring to an oral disclosure, use, exhibition or other means
- *P* document published prior to the international filing date but later than the priority date claimed

- *T* later document published after the International filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
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- *&* document member of the same patent family

Date of the actual completion of the international search

Date of mailing of the international search report

14 May 2002

31/05/2002

Name and mailing address of the ISA
European Patent Office, P.B. 5818 Patentlaan 2
NL - 2280 HV Rijswijk
Tel. (+31-70) 340-2040, Tx. 31 651 epo nl,
Fax: (+31-70) 340-3016

Authorized officer

Herrmann, K

INTERNATIONAL SEARCH REPORT

Inte
nal Application No
PCT/BE 01/00217

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WANG, J ET AL: "Structure and function of microplasminogen: I. Methionine shuffling, chemical proteolysis, and proenzyme activation" PROTEIN SCI, vol. 4, no. 9, September 1995 (1995-09), pages 1758-1767, XP001070157 cited in the application page 1759, right-hand column page 1766, left-hand column -----	40
A	DUMAN JOSEPH G ET AL: "O-mannosylation of Pichia pastoris cellular and recombinant proteins." BIOTECHNOLOGY AND APPLIED BIOCHEMISTRY, vol. 28, no. 1, August 1998 (1998-08), pages 39-45, XP001070866 ISSN: 0885-4513 the whole document -----	1-41
A	CREGG JAMES M ET AL: "Recombinant protein expression in Pichia pastoris." MOLECULAR BIOTECHNOLOGY, vol. 16, no. 1, September 2000 (2000-09), pages 23-52, XP001078868 ISSN: 1073-6085 page 36, line 2-4 -----	1-41

INTERNATIONAL SEARCH REPORT

Information on patent family members

Intel 1st Application No

PCT/BE 01/00217

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WO 9307893	A 29-04-1993	US	5288489	A	22-02-1994
		WO	9307893	A1	29-04-1993
		US	5407673	A	18-04-1995
WO 0018436	A 06-04-2000	EP	1062953	A1	27-12-2000
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		EP	1117437	A1	25-07-2001